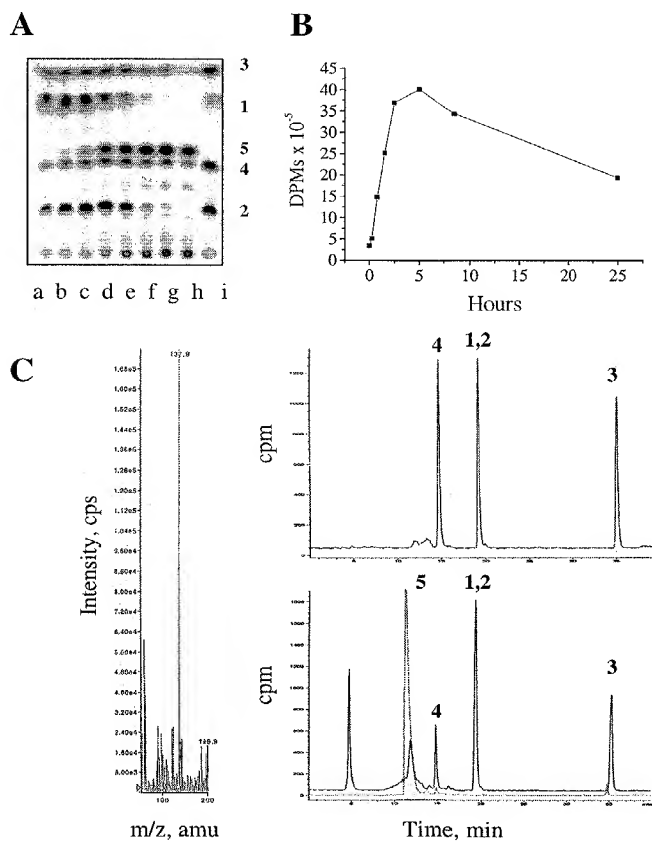
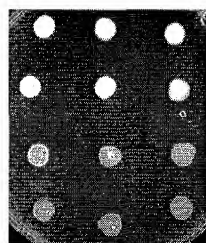


103330" 0363350



Control

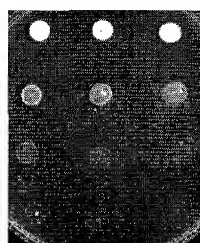


a

b

c

INH 12.5

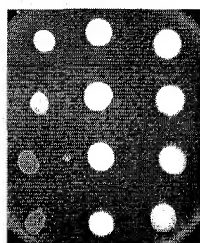


a

b

c

ETA 2.5

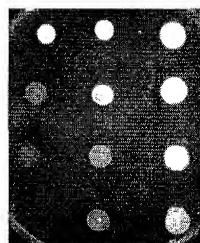


a

b

c

ETA 12.5

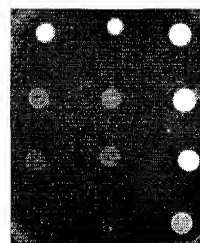


a

b

c

ETA 62.5



a

b

c

FIG. 2

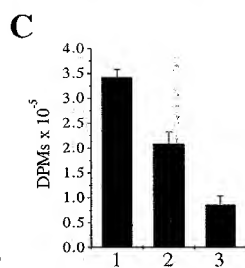
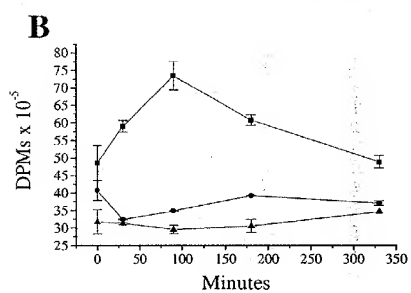
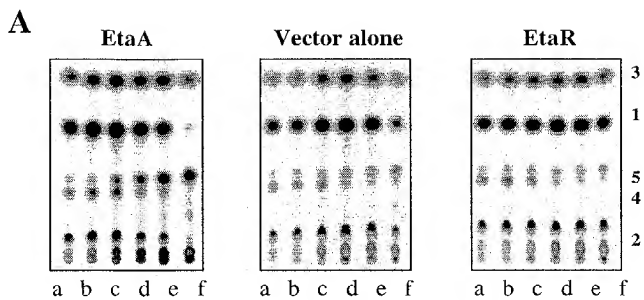


FIG. 3

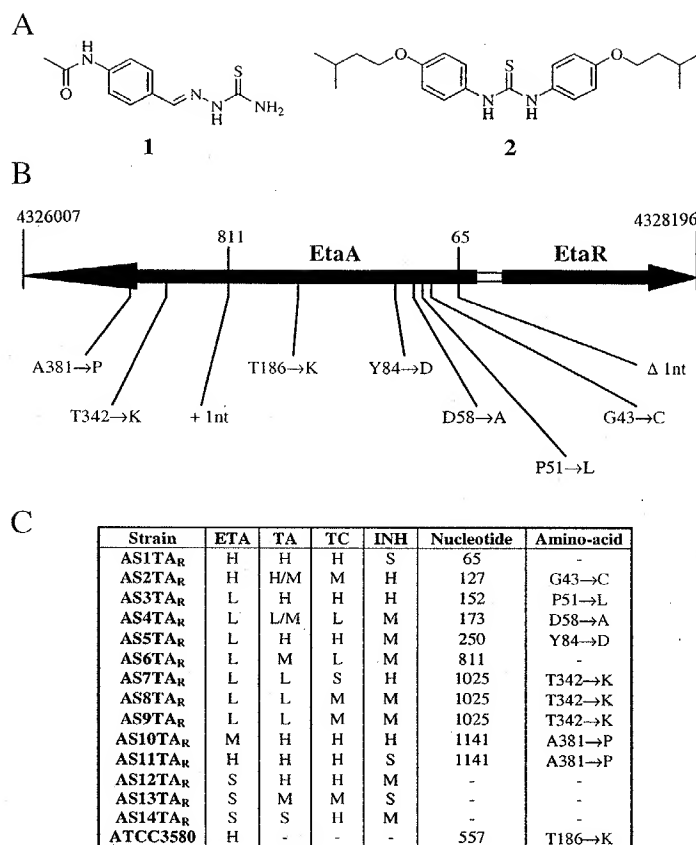


FIG. 4

SEQ ID No.: 1

EtaA: 1467 bp - M. tuberculosis -

acgagg
acgggtcctcgagaaggtttctcggcgggtggcgaggatcgc
cagttcacgatacgtcgccggacggccgcgggtgcgccg
gcccctagggcagcgaagcctgactggccgcggagggtggt
caccctggcagcttactacgtgtcgatagtgtcgacatc
tcggtgaaggcctcgacattacgttgatagcgtggatcc
1 - atg acc gag cac ctc gac gtt gtc atc gtg
31 - ggc gct gga atc tcc ggt gtc agc gcg gcc
61 - tgg cac ctg cag gac cgt tgc ccg acc aag
91 - agc tac gcc atc ctg gaa aag cgg gaa tcc
121 - atg ggc ggc acc tgg gat ttg ttc cgt tat
151 - ccc gga att cgc tcc gac tcc gac atg tac
181 - acg cta ggt ttc cga ttc cgt ccc tgg acc
211 - gga cgg cag gcg atc gcc gac ggc aag ccc
241 - atc ctc gag tac gtc aag agc acc gcg gcc
271 - atg tat gga atc gac agg cat atc cgg ttc
301 - cac cac aag gtg atc agt gcc gat tgg tcg
331 - acc gcg gaa aac cgc tgg acc gtt cac atc
361 - caa agc cac ggc acg ctc agc gcc ctc acc
391 - tgc gaa ttc ctc ttt ctg tgc agc ggc tac
421 - tac aac tac gac gag ggc tac tcg ccg aga
451 - ttc gcc ggc tcg gag gat ttc gtc ggg ccg
481 - atc atc cat ccg cag cac tgg ccc gag gac
511 - ctc gac tac gac gct aag aac atc gtc gtg
541 - atc ggc agt ggc gca acg gcg gtc acg ctc
571 - gtg ccg gcg ctg gcg gac tcg ggc gcc aag
601 - cac gtc acg atg ctg cag cgc tca ccc acc
631 - tac atc gtg tcg cag cca gac cgg gac ggc
661 - atc gcc gag aag ctc aac cgc tgg ctg ccg
691 - gag acc atg gcc tac acc gcg gta cgg tgg
721 - aag aac gtg ctg cgc cag gcg gcc gtg tac
751 - agc gcc tgc cag aag tgg cca cgg cgc atg
781 - cgg aag atg ttc ctg agc ctg atc cag cgc
811 - cag cta ccc gag ggg tac gac gtg cga aag
841 - cac ttc ggc ccg cac tac aac ccc tgg gac
871 - cag cga ttg tgc ttg gtg ccc aac ggc gac
901 - ctg ttc cgg gcc att cgt cac ggg aag gtc
931 - gag gtg gtg acc gac acc att gaa cgg ttc
961 - acc gcg acc gga atc cgg ctg aac tca ggt
991 - cgc gaa ctg ccg gct gac atc atc att acc
1021 - gca acg ggg ttg aac ctg cag ctt ttt ggt
1051 - ggg gcg acg gcg act atc gac gga caa caa
1081 - gtg gac atc acc acg acg atg gcc tac aag
1111 - ggc atg atg ctt tcc ggc atc ccc aac atg
1141 - gcc tac acg gtt ggc tac acc aat gcc tcc
1171 - tgg acg ctg aag gcc gac ctg gtg tcg gag
1201 - ttt gtc tgt cgc ttg ttg aat tac atg gac
1231 - gac aac ggt ttt gac acc gtg gtc gtc gag
1261 - cga ccg ggc tca gat gtc gaa gag cgg ccc
1291 - ttc atg gag ttc acc cca ggt tac gtg ctg
1321 - cgc tcg ctg gac gag ctg ccc aag cag ggt
1351 - tcg cgt aca ccg tgg cgc ctg aat cag aac
1381 - tac cta cgt gac atc cgg ctc atc cgg cgc
1411 - ggc aag atc gac gac gag ggt ctg cgg ttc
1441 - gcc aaa agg cct gcc ccg gtg ggg gtt
tagcttttagcgacggttttagcgccggtttaggccatagt
cagacgacgatgatgccgtcgctcgctcgctgtaggcgata
tcgcccgggaacgaatgtcaccocgccagcgtgatttca
acgtcgcggtttctccggcaccgggtcttgggtgctcttgagg
ggattgggtgccagcgctttgatgccgatgtcgatgccg
cgacg

Figure 5

SEQ ID No.: 2

EtaA: 489 aa - M. tuberculosis -

1 - MTEHLDVVIV GAGISGVSAW WHLQDRCPTK SYAILEKRES MGGTWDLFRY PGIRSDSDMY
61 - TLGFRFRPWT GRQAIADGKP ILEYVKSTAA MYGIDRHIF HHKVISADWS TAENRWTVHI
121 - QSHGTLTALT CEFLFLCSGY YNYDEGYSPR FAGSEDFVGP IIHPQHWPE LDYDAKNIVV
181 - IGS GATAVTL VPALADSGAK HVTMLQRSPT YIVSQPDRDG IAEKLNRLWP ETMAYTAVRW
241 - KNLVRQAAYV SACQKWPRRM RKMFLSLIQR QLPEGYDVRK HFGPHYNPWD QRLCLVPNGD
301 - LFRAIRHGKV EVVTDTIERF TATGIRLNSG RELPADIIIT ATGLNLQLFG GATATIDGQQ
361 - VDITTTMAYK GMMLSGIPNM AYTVGYNAS WTLKADLVSE FVCRLNLYMD DNGFDTVVVE
421 - RPGSDVEERP FMEFTPGYVL RSLDELPKQG SRTPWRLNQY YLRDIRLIRR GKIDDEGLRF
481 - AKRPAPVGV

Figure 6